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<110> Sun, Yongming Recipon, Herve Ghosh, Malavika Liu. Chenghua

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| acttttggaa | gagacatttn | taactagtaa | tgataatttg | tcttaattca | taaacacttc | 5820 |
| aaatcacata | actgaataca | ttttcaacca | ggaggatgca | acattacccc | aaaataccga | 5880 |
| gtcanagaaa | ttattattgt | tgagacaacc | aggtaccaaa | ctcttaattc | ccattgggtt | 5940 |
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| tactttcagg | ggtatcctag | taaaagcgac | tttccatgtg | cctgcctgtg | cctttctgtg | 120 |
| ccagggtctt | aagaaatgtg | tgtgggtttc | ctctgtggct | ctgacactcc | atcaaacacc | 180 |
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| gcttaggtgc | tgctttaaaa | aataagtagc | agtgtattcc | agactcctct | aagaaaagag | 360 |
| aattgctcat | tgtggaacag | gcatggcaat | gcagtgcctt | tgcctggaac | accttgctgc | 420 |
| tgccatcact | aagacccatt | ctgggacaag | aggaggctta | ccccttattg | agtatctgcc | 480 |
| atgagetttg | tgtggtcctt | gnnnnnnnnn | nnnnnnnnn | nnnnnnnnn | nnnnnnnnn | 540 |
| nnnnnnnnn | nnnnnnnnn | nnnnnggcca | cccatttcct | gacagagatt | ggcctacaaa | 600 |
| ggtcagatgg | cttccagggc | acactgagcg | gcccttgtgt | gtcggaatgt | tecettetgt | 660 |
| ttgtcccttc | caggctggac | actttgggag | cagaagtcaa | agacaccttt | atcattgtac | 720 |
| cctcagcacc | tggtgtagtg | cctgggattt | agtagttctg | aggagcgtgt | gttgaatgaa | 780 |
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| ttttctattt | attatttaat | actttcaggg | gtatcctagt | aaaagcgact | ttccatgtgc | 120 |
| ctgcctgtgc | ctttctgtgc | cagggtctta | agaaatgtgt | gtgggtttcc | tctgtggctc | 180 |
| tgacactcca | tcaaacacca | gggcttgtgc | acgcaggtga | gaagcatacc | caggtgcctc | 240 |
| agagatattg | aagagggtct | ggaatgtggg | agagaggcaa | atggctttcc | aataaaagta | 300 |
| ggctaagaca | taattagggg | cttaggtgct | gctttaaaaa | ataagtagca | gtgtattcca | 360 |
| gactcctcta | agaaaagaga | attgctcatt | gtggaacagg | catggcaatg | cagtgccttt | 420 |
| gcctggaaca | ccttgctgct | gccatcacta | agacccattc | tgggacaaga | ggaggcttac | 480 |
| cccttattga | gtatctgcca | tgagctttgt | gtggtccttg | cttgctaggt | acttgagcaa | 540 |
| gcattatctt | ttttaccttt | taaaacacca | ctgaggtgta | ggtatggcac | ccatttcctg | 600 |
| acagagattg | gcctacaaag | gtcagatggc | ttccagggca | cactgagcgg | cccttgtgtg | 660 |
| teggaatgtt | cccttctgtt | tgtcccttcc | aggctggaca | ctttgggagc | agaagtcaaa | 720 |
| gacaccttta | tcattgtacc | ctcagcacct | ggtgtagtgc | ctgggattta | gtagttctga | 780 |
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ccgtctgtag catgggaggg gcctggagca tagcagggcc tctcacgggc tttgntttca
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ggttgacatt t
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                                                                     120
                                                                     180
aatggtgtga totoggotoa otgoaacoto cacotocoag gotoaagtga ttotoctgoo
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tttttagtag agatagggtt teaceatgtt ggeeaggetg gteeteaact eetgaettea
                                                                     300
                                                                     360
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180
                                                                     240
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aggaagtgac tagtttggtt aagagctggg aactgagtca ggtaagccgt gtcatgttgt
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| aactccacca gaaaatggag gagagcgggt ttccaggaga caaagctgag atga | agaagtg 240 |
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| aggettgett geetacteet tetetettte eagagggaaa eettgtggtg gtte | ectcact 360 |
| gtotattoat tatgcaagga aatgagggot tttaagggtt cotcagattt ttot | ccacca 420 |
| aagagtgett teacaagtta ttgaggegtt tgttteeatt ttaaagtaaa ettt | tggaat 480 |
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| agtcaccagg aaatgtgatt teeteettgt gaagatggtg atggeeetaa gete | gagattt 660 |
| ttttgagttc tagggtttgg ttatcatcat gttttgatgc attgcaagac ttta | attgtct 720 |
| gatttgagtt gatttctgca aatataaaaa ataataagaa taatcctgca agat | ctcaga 780 |
| ggaactctaa gactggctaa caccagtttc tccaggttct ccatttctct tcag | ggtcgtt 840 |
| ccatttgtat gttaggcctt ctttcagttt ctttgtttcc cctttccctt tccg | gtegget 900 |
| aattttttct gtgttctgaa gtactcttaa gtcttcagaa atatcagtat gtct | tcttaa 960 |
| caatgtcgct atggaaacaa attttaaaaa catgatgtca gttgagaaaa cctt | atgtcc 1020 |
| aggtatette acetttttaa ttgggaggaa tttattaate atgtaggaag acat | tttgtg 1080 |
| aggataattt gaaaaaagga cccagtgcta ccctagtcca cacacattga tggg | gagetet 1140 |
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| aatttgtoot ggttttotag tacotcaagg cagatatgca aaggtgttta ggag | gacatac 180 |
| totoagacaa accattatta ttttaaagga tagaacaaaa caatogotag ttaa | iggaaga 240 |
| tgttttgtaa taattaaact tgtaattatt tgacttgaaa tatttaatca tttt | tttggg 300 |
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| ccgtgtctcc catattctcc ttgtgtttga aacataaaac aaacactaaa ccta | |
| agttgctggg tttgttttca taattgaggt gagtttttcc ctcaactatt acaa | itaaaag 480 |
| aaaacttttt atgattttaa tgataatgtt ttgtggtggg ttaaagacct ccta | lacaaca 540 |
| | |

| gggggttttt atacaacaac | aagaagtttt | taaataattg | agtttttaaa | gtggaaagca | 600 |
|---|------------|------------|------------|------------|------|
| gcagtaaatt aaactagaag | gatatatttt | atacctagaa | ataaataaag | ctcaacttgt | 660 |
| tttgtaagcc tgttttaaaa | atatttaatc | atttaatttg | tgcaagtata | gagttctcct | 720 |
| atggcaaaac tataccatca | tcttctccaa | ttgtgcatgg | cagctgtact | aagttctgca | 780 |
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| gcagttcaag caattctact | gcctcagcct | ctcaagtagc | tgggactata | gacattcacc | 180 |
| accacaccca gctaattttt | tgtattttta | gaaaattttg | tatatttaga | aaaggtttca | 240 |
| ccatgttggc caggctggtc | ttgcactcct | gacctcaggt | gateegeeca | cctcagcctc | 300 |
| ccaaagtgct gggattacag | gcctgagcca | ctgtgcccag | ccctcaagta | actcttaaac | 360 |
| ctactgaagt tagacaatca | ataactgaaa | tgacatcatc | tttcttgaat | gtttaaggaa | 420 |
| ataaagttcc ttcttctgac | aaactttaag | tgtgttcttg | atttccttgc | ctccctcttc | 480 |
| ctctgggagt tttcttccct | agctgctcac | tttcattatc | aacgaaatat | tcctcttcac | 540 |
| gcctttttac cttataccta | caacatgctc | agttctctct | ctttacaaga | aaatataagt | 600 |
| tttcaccaac ctatttatca | aatttacatc | cccctccctt | tctacttcct | tttgtaaaaa | 660 |
| aagagcattc aacctattgt | ctgtctccat | gccctcacat | tatcagtgca | agcacccgca | 720 |
| actgtggctc tccaccatgt | gageteaace | tatcatcaca | actgtatctc | ccctaacact | 780 |
| catttagatt aagccatttt | tcacaagttt | ctaaaattat | ctcttccatt | tctcagtata | 840 |
| accettett teeetteaca | gtttcttgaa | ccaatctcac | tagtccttca | acgttcactt | 900 |
| ccaaggccac cccgaacaca | tcttttcctc | ttccctaaat | aaattctact | ggattctttc | 960 |
| tgtttttcac tggaaacttc | tcatactcca | ttggttcctt | tctcatgaca | tttattttac | 1020 |
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| agaactggac cagtgtctct | tatatttata | tccctaataa | tacttattaa | acacgtagtc | 1140 |
| tattctcaac attgaattcc | atcttatact | caaagaataa | tactttaaca | tagccattgt | 1200 |
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| | | | 24 | | | |
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| ttatcttcag | ttatgctata | gcatgtacat | ttccattctc | ttgtcgaagt | ttctttcgtt | 1380 |
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| gggggttcca | cttcatgagt | agtgatagac | cagcaatcac | tatacttgac | actaaaccta | 1560 |
| aacctggcta | taaaatatta | ccaatttcta | agggggtatt | tatgttgact | gtatataaat | 1620 |
| ccatttccag | agggcttata | tttaaatgtg | tcttgatata | ccaattctga | gaaggtatga | 1680 |
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| agatgatggt | atagttttgc | cataggagaa | ctctatactt | gcacaaatta | aatgattaaa | 1800 |
| tatttttaaa a | acaggettae | aaaacaagtt | gagctttatt | tatttctagg | tataaaatat | 1860 |
| atccttctag | tttaatttac | tgctgctttc | cactttaaaa | actcaattat | ttaaaaactt | 1920 |
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| ttatgaaaac | aaacccagca | acttttgctt | aggtttagtg | tttgttttat | gtttcaaaca | 2100 |
| caaggagaat a | atgggagaca | cggggaaaaa | agccactgct | taattttaag | agtgctaaca | 2160 |
| ttaacaaaat (| ctatccattc | tttcccaaaa | aaatgattaa | atatttcaag | tcaaataatt | 2220 |
| acaagtttaa · | ttattacaaa | acatcttcct | taactagcga | ttgttttgtt | ctatccttta | 2280 |
| aaataataat 🤉 | ggtttgtctg | agagtatgtc | tcctaaacac | ctttgcatat | ctgccttgag | 2340 |
| gtactagaaa a | accaggacaa | attctagtgt | gtgcaaaata | aatttaagct | acatatcaaa | 2400 |
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| cttagcagtg gttattgcat | agaacaactc | cttacacaga | gatttgcaag | ctttctgaac | 420 |
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| agattagtat attaacaact | aataacaata | ttaaaagtta | gaacaattcc | tttcctctat | 540 |
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| tctttcattt | tcactttctt | ccttacactt | gcaatccaga | gtccagatgt | aaaacagtgt | 180 |
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| gtgtccttgg | gatggtaccc | ttacccctga | ggtgctaggg | atgggcccca | gggtctttcc | 300 |
| ctgctttcta | ctttcctaat | ggctaagtga | tgtcagagga | caacatcttg | atgtgtagag | 360 |
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| ccaattgttt | tcaggttttt | tactaggagc | acatgcatga | atgtgtatat | atgtgtatag | 480 |
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| taatgttaac | cacatgctat | atacttatat | ttttctttca | tttgcaaaag | aatgctgtta | 660 |
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| geeteteett | ggcttgcaag | atggccacct | tctggctgtg | tcctctctct | catggccttt | 180 |
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| aaacagagag cttccaggag | gatcaatgcc | attcaatgag | cttgctgctg | tactcccctc | 180 |
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| agaccgaaac agagagcttc | caggaggatc | aatgccattc | aatgagcttg | ctgctgtact | 180 |
| cccctctaca caatatggat | atatcccatc | ccagcccgag | actggccata | ctagttctag | 240 |
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| atacaataga aagatcctgg | aatcccgaca | tgaggacaaa | aatggtactg | aattcttttt | 180 |
| gaaaaataga ttactgaaaa | gcgatctaat | atagaacagt | tgcttttact | tagatgttca | 240 |
| atgcatattt gttgtataat | aaccaagtta | ttacagttca | gataaagggt | ccaaagtgtt | 300 |

| ttcgttatga tataatactt tctattgtaa actggactaa agaaacgtt | ; tatgttcaag | 360 |
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| gcaggagtca cagacetgca ggcaegeact tgccagtgae tgggaegtt | g gctggtggtt | 480 |
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| taatgetaet gaacagetae agageaetee tetgaaetea etggaatgg | g ctatatccca | 180 |
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| | gaatacagac | | | | | 180 |
| | ttcaaatgtg | | | | | 240 |
| | cactcacgag | | | | | 300 |
| | gctggggtct | | | | | 360 |
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| toatgaaatg aatgootttt tacttgaaag aatgactgag agccaggota tggatattca | 180 |

| aacatgtatt tttcagacac ttcttgaaaa taagtgaagc aaacctgtta attacaaggg | 240 |
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| atcatttcat gtgtccactc tttttaaaaa tgttaccttt tctaggattg gcagaatttg | 300 |
| gaattatatg tottattaaa tatgotttga aagacagaag taataagtto tggttaatot | 360 |
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| aatctaacat atcaaaatta gagaacccca aaccatcaca ttetttetet ttgtgccatt | 480 |
| ttagaattga gaataccgtc cttcttactg tggttatatt tttacttttg tatataaact | 540 |
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| 210 50 | |
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| cccgctcagc aacaataaca acctgcattt agggaacgtg tgttatgtgc caggccacac | 180 |
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| tccaaaccte agagacagee eccatecete tegttageea eccaeacace eegeteagea | 180 |
| acaataacaa cetgeattta gggaaegtgt gttatgtgee aggeeacaea ggeattatet | 240 |
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| agcagttete cetacetagg cetttettge teaccatg | ta aaagattoot atttagttto 420 |
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| aggaagcaat tgcatatcaa ttttcttcta ttccagct | |
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| aagaaggaag caattgcata tcaattttct tctattcc | ag cttaatctat ttatttttct 180 |
| cttttacatt aaaacattct tttaatgata tatgctgc | ct gtaaatattt cccacccact 240 |
| ttccagaggt aatccactgt tatcaagtaa gtttagta | aa ttttttttaa ttgaattttc 300 |
| tcaataggtc attaacgtgt ttcaaagttg aaaaatta | ca aaactatgtg tcgtgaaaag 360 |
| teteettett teeettgtgt eecaagetae etagttet | tg gagccagttg atgttatcag 420 |
| attettttgt attettteag acacacatgg tatgeatt | tt tgagcaaagg ggcgtgggtg 480 |

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| tttcttaagt aactttattt cattatttgt attcagtttt gtaaaattag atactacatg | 600 |
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| gcactgaatg aggaccagca ggaagagatc tcagaaaaca taagataatg gacttgttga | 360 |
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| gaggaaacta tacaatgacc tagcaaattt gaaaaaggaa ccaggaacaa cttgtagaca | 480 |
| tgaaaagttc atgtctttat aataaaaatc taacagatgg atttactagc agattacata | 540 |
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| nnnnntaact ctaatacacc | agtacaatgg | aaagatteet | aaattcaaaa | gccagaaggc | 180 |
| tgggttcctg ttcccaccct | gccttttacc | ttctgtgtgt | tcctgatgaa | gacacttcat | 240 |
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| tgcttggagg gactcagagg | ctgtggagag | tgaatatttc | catcagctga | tgcccttctc | 360 |
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| gcaggctgtc agccacgggc | aagctttctg | aaaaagacgt | gccaactgca | gccacagaaa | 540 |
| gtccattccc ttgaataact | ctgctaatat | ttgaaaatta | gttcccttgc | tcctgatcat | 600 |
| gctactgggt atttggatat | aagagccaag | gatgagggca | atagaaaatt | aaaatcatgt | 660 |
| tctactcata taaactgcac | agatatggaa | gggtaggtcc | tattacctat | aatcctggga | 720 |
| tttttagact ctcactttca | ttggaccaga | gttgccttag | ggacagtaaa | aacacaaaat | 780 |
| gctgggtatt gttttcatca | agcaactact | gatagtgcac | atttaaatca | aaattcttct | 840 |
| aatcccaaac tcagtaaaca | gatgctgtga | gcttagttct | gcccctctgg | cttcagattt | 900 |
| taccccactg gatgtgccca | attctgagat | gacaagacgc | ttccagcttc | cacatggttg | 960 |
| caatttggct gtggaactgg | catgaaagca | cgtcactgtg | tcagcacctg | ggccaccaga | 1020 |
| tgaataacct atgaacaaca | gctttggact | aaaatatgaa | ggggttgttt | tccttcaatc | 1080 |
| tececetace ttectcagaa | cctgctacaa | ggaaagattt | atagactcga | aagcgtcaat | 1140 |
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| aaangnataa aaataa | | | | | 1216 |
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| ggcctccagc ttgcagatgg | catatcatgg | aacttctcag | cctccaaatt | cataactcta | 240 |
| atacaccagt acaatggaaa | gattcctaaa | ttcaaaagcc | agaaggctgg | gttcctgttc | 300 |
| ccaccctgcc ttttaccttc | tgtgtgttcc | tgatgaagac | acttcatgct | ccactattta | 360 |

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| tcagaggctg | tggagagtga | atatttccat | cagctgatgc | ccttctcagg | cctcaatctc | 480 |
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| tcagccctgt | tttccatcac | caaaggactc | cggaggaact | gtgccaagca | ggctgtcagc | 600 |
| cacgggcaag | ctttctgaaa | aagacgtgcc | aactgcagcc | acagaaagtc | cattcccttg | 660 |
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| ttcatcaagc | aactactgat | agtgcacatt | taaatcaaaa | ttcttctaat | cccaaactca | 960 |
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| gaactggcat | gaaagcacgt | cactgtgtca | gcacctgggc | caccagatga | ataacctatg | 1140 |
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| nnnnnnnnn nnnnnnnnn nnnnnnnnn nnnnnnnn | 300 |
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| cttttaagga tatagagaaa gcaaaattag caaatctagt ttcttgtcac tttactagga | |
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| atgccagtta gagcgttaat gccacacgag ccagagaggt caccttg | ctg agcatggctt 360 |
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| a | 541 |
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| agateetaae agagtgeate ttgtgetttt eetaaeagae etgtegge | act ggctttttct 180 |
| cttttaagga tatagagaaa gcaaaattag caaatctagt ttcttgt | cac tttactagga 240 |
| gggaggaaaa gagagaaaga atgcacttgg gaatgggagg ccttgct | ttt aatttaccag 300 |
| atgccagtta gagcgttaat gccacacgag ccagagaggt caccttg | ctg agcatggctt 360 |
| gactgttgca gcctctttct gcgactccag acatgcgatg tctgttag | get gattetagee 420 |
| ttcagatgca gcccggagat gtaaccctga ggctggagtc ctgtggc | ct aatcccagac 480 |
| agaggcaact ccaccaagtt ctggtttggg tcagaaatag agggaaa | gga tgaatgaaag 540 |
| aagatacaaa gaaataatga acaagtgagt tettteaget gettaett | tgg gtggtctgca 600 |
| ggcagcaaga gacaggaagg aggctgttgt ggggtccttg ttcgagg | cag tgggagattt 660 |
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| geocaggeet geaccectaa ggeaggeact geteegtgat ceaggaac | |
| cagetgggag tgageagtea gagagggaga cageettgee eggtgeta | |
| gtcaccgagt gggcagaggg aggageggcc ctcaccggat gtcaagca | age etgggteece 240 |

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agtocagete tgeetgteee tegeaataae geeteagtga egaceatttg tgagecatet
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ctctgtctca ggcacggtgc tacatgccaa cgaaacctgc tcccattgaa ccctggccag
                                                                      360
ccagtgaaga aagggttggg cctgggaggt gccactttac agacaggggc accaaggggc
                                                                      420
agggtggcag gaggcccacc ggacgttccc catgaagtag cagtcccagc atccacaccc
                                                                      480
                                                                      540
agcaggeace acgetggece geagectoce tgccagcacg cetggettee eggectegga
acttgatctg ctccctcttc cggacactgg ggctcctgcc aagtcctggg ctgggcagca
                                                                      600
                                                                      660
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tggccggcac ctacttgtgg ggctgggggt tcccccagca ggttgggctc cccacctgac
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                                                                      900
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ccaggggctg ggcctgtccc agctctggtc ccccggcccc aggtcctgga cgctggntcc
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agatgagtet gegeeggate gacetgetge egagteetge eggacaggea eaggeaggga
                                                                     1080
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cattttactt tacctgtcta cagtgttttg cgcaattgac cactccttcc tttttgaagt
                                                                                120
attitiette ettggtitet gaaataetgi tatetteeta teteaetgge catacattet
                                                                                180
agteteettt getagtttat tatggtttte atetteteaa caacaatttt nttttttng
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ctcttagtaa ttagttttgt tttgttttgt tttgttttaa tgttgtgctt atcttaaggt
                                                                              120
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| gtaaatgcag acaaagttgg | aattgaagct | gccgaaatgc | tattagcaaa | tcttagacat | 180 |
|---|------------|------------|------------|------------|-----|
| ggtggtactg tggatgagta | tctgcaagac | caggtaatga | cacatttagg | ttaaaaaccc | 240 |
| tctaacctgt tagatttgaa | tatgtggtag | attgaatatc | aatttaaata | attgactttc | 300 |
| agacactaat tagcaagtcc | tacttcaata | atttaaaaaa | atattctggg | atttgcattc | 360 |
| ctcaaatttc agccctcatt | ttactttacc | tgtctacagt | gttttgcgca | attgaccact | 420 |
| ccttcctttt tgaagtattt | tctttccttg | gtttctgaaa | tactgttatc | ttcctatctc | 480 |
| actggccata cattctagtc | tcctttgcta | gtttattatg | gttttcatct | tctcaacaac | 540 |
| aattttnttt ttttnggngg | aganggagtc | ttgcnatgt | | | 579 |
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| ccttgaaagc tggaaagagc | ttaacaaata | tcagctgttg | ccatgaaaga | atatttgctt | 120 |
| actttccatt gtgtataaga | taacgataat | catagaatta | atattattca | acttccttgt | 180 |
| gtcttttgca catttctgta | cagtcctgtt | tttgtttgtt | actgtcattc | tcaaagtact | 240 |
| caagttgaat tttgtcactt | tggatttctt | ccaggaatat | gtgagagaca | tttaggtctc | 300 |
| taatgatgaa gtattttcta | ggcgtaatgc | aaaagattg | | | 339 |
| <210> 75 <211> 299 <212> DNA <213> Homo sapiens | | | | | |
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| ggtgacttcg tttccacctc | cccttatata | ttgttcttcc | ttcctctcta | aattctctaa | 120 |
| atctctgctt atacagagca | atctggctct | ctctggcctc | tccagtcatc | atacatcata | 180 |
| ctcacattca ccatcttgag | aagtgcagta | agccacataa | atgcagcaga | agtaccttat | 240 |
| gcagtcctag gaggctgtgg | ttttgagttg | ctttttttt | tcttttggga | gacggagcc | 299 |
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gaaatggagg ctcagaggga tatgtagtag ctaaatgtta gagctaggat tganacccaa 180
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<210> 77 <211> 254 <212> DNA <213> Homo sapiens

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tgaaatggag gctcagaggg atatgtagta gctaaatgtt agagctagga ttgaaaccca 180
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gtctttacat ggcgttcccn nnnnnnnnn nnnnnnnnc aaatttcctc ttttcataag
gaccgtggta ttggataggg gtccacccta cttcgatatg accttatttt aantncatct
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gaaggtggaa aggggacctg agtccaggca tgtgggcagc ctggagaagg cgaganaatg
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gattettece cagaateeet ggaaaggnae gtggeeetaa e
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atateteatt geataetttt atgtaaettt gtettagaaa aacaagagtt et
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<212> DNA
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gacaatttaa aaacctatac aaagagtgac acataaataa acaaaaacaa cataaaaaata
aaaatataat totaaaaata ttoaagtago caattggaag gtggaaaaaa gaaaaagaac
                                                                      180
aaaaaataga acagcactaa acaaaaaata aaatcgcaga cctaggccct gacatatcaa
                                                                      240
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taattatatt aacatgtaaa tggtctaaat tttacca
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aactcagatg tattaatttc ctattgtgtc tnnnnnnnn nnnnnnnnn nnnnnnnnn
                                                                     120
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| nnnnnnnnn | nnnnnnnnn | nnnnnnnnn | nnnnnnnnn | nnnnnnnnn | nnnnnnnnn | 180 |
|-------------------------|------------|------------|------------|------------|------------|-----|
| nnnnnnnnn | nnnnnncact | ttctttctgt | aggctctagg | agagaatcta | gnnnnnnnn | 240 |
| nnnnnnnnn | nnnnnnnnn | nnnnnnnnn | nnnnnnnnn | nnnnnnnnn | nnnnnnnnn | 300 |
| nnnnnnnnnn | nnnnnnccaa | gtccttctca | cactgctgtc | tttttggttc | tctctcttgc | 360 |
| ctgcctnnnn | nnnnnnnnn | nnnnnnnnn | nnnnnnnnn | nnnnnnnnn | nnnnnnnnn | 420 |
| nnnnnnncat | agttgattag | cagccttaat | ccatctgtaa | ttttaattcc | cttttgccag | 480 |
| gtaatgtggc | cattatcttg | cctacaacct | cagaggatgt | tgataatgta | aagggtagtg | 540 |
| aattggggag | ttcatagggt | ttgatagttg | acaaatacag | agtgtagtat | taggtagggg | 600 |
| ttttttggca | gggtgcagtg | gcccatacct | gtaatgt | | | 637 |
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| tggagaaaaa | tttaataatc | ctggcagggc | tacattcaac | ataattctgt | tatgggggaa | 120 |
| ggcagcatgc | tttggctgct | cagtgagcta | tgttctgtac | aaccaagtga | aattgctaaa | 180 |
| aaaagattct | cctgtataca | gtaacttaaa | gtgatgcagt | ctacttaaga | tcagatctga | 240 |
| gttacaaaat | caaaagtgac | agctcctatg | ttcttttaaa | gtccaatctc | tttttttcat | 300 |
| tgttgtgctc | caaatgcctt | gagtacctga | tgtagagtag | gtggctaata | aatattggtt | 360 |
| gaatttettg | aacgaatctg | ttatgaaaag | atctactttg | ctcatctctg | tgccccaata | 420 |
| gcaggagctt | gaggagaagg | agaaaatatt | gggtcagagc | ttttgattaa | tatgtatgat | 480 |
| tctattaaac | gggttcacta | aaccaaaaaa | ggcaaggaaa | acagttaaac | caagagtctt | 540 |
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| gctcagggct | catggctgat | attacaggca | taagccacca | cacctagcca | agaaaccatt | 120 |
| ctttgaacac | aagcaaatat | actttggaga | aaaatttaat | aatcctggca | gggctacatt | 180 |
| caacataatt | ctgttatggg | ggaaggcagc | atgctttggc | tgctcagtga | gctatgttct | 240 |

| gtacaaccaa gtgaaattgc | taaaaaaaga | ttctcctgta | tacagtaact | taaagtgatg | 300 |
|--|------------|------------|------------|------------|-----|
| cagtctactt aagatcagat | ctgagttaca | aaatcaaaag | tgacagetee | tatgttcttt | 360 |
| taaagtccaa tctcttttt | tcattgttgt | gctccaaatg | ccttgagtac | ctgatgtaga | 420 |
| gtaggtggct aataaatatt | ggttgaattt | cttgaacgaa | tctgttatga | aaagatctac | 480 |
| tttgctcatc tctgtgcccc | aatagcagga | gcttgaggag | aaggagaaaa | tattgggtca | 540 |
| gagcttttga ttaatatgta | tgattctatt | aaacgggttc | actaaaccaa | aaaaggcaaa | 600 |
| ggaaaacagt taaaccaaga | gttcttgagg | ttaaagtett | gtgatgatta | aaatcatcat | 660 |
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| cttgcatgca catcccc | | | | | 77 |
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| atgccaaagc tggtataaga | caccaaaaga | gtaagacaga | aagtattctt | ccctggagct | 120 |
| ttgtctgact ttccaagctt | tattaggcat | caaacaaaac | tgaagtgctt | tttaagattc | 180 |
| aagteteeta egtegtetaa | ggcagagtaa | gtagccttca | gtactatatt | ttactctaat | 240 |
| ttttttttaa cacaatggca | gtactataag | tatgaaactt | tggtataaat | gtcagattct | 300 |
| agattgtgct cctgctttct | gcacactcta | atatttttaa | acatctcgaa | aatacagagt | 360 |
| ggcagcaaaa ttacctgtaa | aaacatacta | gctcaagagt | ttgacaggct | caaaataaat | 420 |
| taccttaaat acattaaaca | agaagtgtat | ttgttataca | gtatgtactg | accaaaatta | 480 |
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| aagtgcaggt tgtacagaaa | gagctgcttg | tgttatttta | tgagcaaaat | gaaaagctaa | 540 |

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<400> 88

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| gccccaccct actcccagc | a tatgcacaca | cacacgtgca | cacacaatac | tcacttaaca | 120 |
| aacatttaat ttattgaac | a tttattatat | gccaaagctg | gtataagaca | ccaaaagagt | 180 |
| aagacagaaa gtattcttc | c ctggagcttt | gtctgacttt | ccaagcttta | ttaggcatca | 240 |
| aacaaaactg aagtgcttt | t taagattcaa | gtctcctacg | tegtetaagg | cagagtaagt | 300 |
| agccttcagt actatattt | t actctaattt | ttttttaaca | caatggcagt | actataagta | 360 |
| tgaaactttg gtataaatg | t cagattctag | attgtgctcc | tgctttctgc | acactctaat | 420 |
| atttttaaac atctcgaaa | a tacagagtgg | cagcaaaatt | acctgtaaaa | acatactagc | 480 |
| tcaagagttt gacaggctc | a aaataaatta | ccttaaatac | attaaacaag | aagtgtattt | 540 |
| gttatacagt atgtactga | c caaaattaaa | gtgcaggttg | tacagaaaga | gctgcttgtg | 600 |
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| aattacaagc gtgacgcac | | | | | 180 |
| | | | | | 240 |
| gtaaaatatt atttgaata | | | | | |
| gaaaaaggct gggtgtggt | | | | | 300 |
| ggcggatccc ttgagctca | | | | | 360 |
| acaaaaatga gccaggcat | | | | | 420 |
| gtgggaggaa ggcttgggc | c taggaggtgg | aggttgcagt | gagccaggat | tgtgccactg | 480 |
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ageetttata gatttgeeat gateetaata catataagea tteattgtat teattattaa
                                                                     180
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                                                                     240
                                                                     300
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                                                                     120
ageetttata gatttgeeat gateetaata eatataagea tteattgtat teattattaa
                                                                     180
ttacttcata gattcagtgt gtgacgaagg gagatgattt ttaacaaata ataaagtgaa
                                                                     240
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tttatagtga tttttttttc actatggtat tttcttaaat atattaagtg cttttcattt
                                                                     360
tetgatacea eetagtttaa ttgggggtga atateagaga aattagaatg ttattteage
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tgaaggagta cagttttttt tttctcttct tagagaatat agtgcctcag atacagtcca
                                                                     480
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nnatattaat nagatgtatc aaactgtnac aatatgtgaa gagtattgtg tatatacaaa
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caggaaacaa ttgaaagcct tcaacatgtg tgggtggggg gagagataac tgaattaaca
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ggccatgtag taaaacttaa aatcaaatcc agtagtcttg aaggtatagt aattgtttag
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                                                                           180
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caggaaacaa ttgaaagcct tcaacatgtg tgggtggggg gagagataac tgaattaaca
                                                                           300
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                                                                           360
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eggtggetea egeetgtaat eecageaett tgggaggeeg aggegggegg ateaeetgag
                                                                           420
                                                                           480
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                                                                     780
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                                                                    1260
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                                                                    1380
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gcccagtgag ggaagtgcaa caagtatggt tctcctgctt gtcaactgtg gaaacagcga
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| tgtttagatt | ttggtactgc | aactgctttc | ctcttgccca | gaaatgtttt | gcctcttctt | 180 |
|---|----------------|------------|------------|------------|------------|-----|
| ttcctacaag | ttaaatgttc | taaatataaa | ggggtatgtg | tgtgtgtgtg | taattctaat | 240 |
| gtgaaaggca | ctagctgtct | aataggtttc | atgtatcatt | actattacta | tatgtatctt | 300 |
| aatgtagtct | atgtaggttt | ttatcagaaa | gtgtaccttt | ctatggttta | ttattttata | 360 |
| ttctggggcc | ttttatctca | gatataaacc | atgaacagta | atgatagtcc | ctgacatata | 420 |
| aatcttagta | aaaagtgatt | aaaaatctaa | aactcagtat | gaaaaacata | tcttgttagc | 480 |
| ataaattaaa | accttttatt | gtttaaaaaa | aaaaaa | | | 516 |
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| acgtgatgtc | actgacccag | gtcattcctt | tactttcaaa | tcttcaagaa | ttggatttat | 180 |
| cagccaacaa | aaagatgggc | agttcttctg | aaaacttact | cagcaggctc | cgatttttac | 240 |
| cagcattgaa | gtcattagtt | atcaacaact | gtgctttgga | gagtgagact | tttacagctc | 300 |
| ttgctgaagc | ctctgttcac | ctctctgctc | tggaagtatt | caacctttct | tgggaacaag | 360 |
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| caaggaagaa • | aacttcctca | gaaggagaaa | ctaagcccca | gacttcaact | gtcaacaaat | 180 |
| ttctcagggg | ctccaatgct | gaaagcagaa | aagaggacaa | tgaccttaaa | acaagtgatt | 240 |
| cccaacccag | cgactggata | cagaagacag | ccacctcaga | gactgctaag | cctctcagtt | 300 |
| cagaaatgga | atggagatcc | agtatggaga | aaaatgagca | tttcctgcag | aagctgggca | 360 |
| aaaaggctgt | caacaagtgt | ctagatttga | ataactgtgg | attaacaaca | gcggacatga | 420 |
| aagaaatggg | agaagcattt | gagatgattc | ctgaacttga | agagctaaat | ttgtcttgga | 480 |
| acagtaaagt | gggaggaaat | ttgcctctga | tccttcagaa | gttccaaaaa | gggagcaaga | 540 |
| tacaaatgat | tgagcttgtg | gcttgctccc | tcacgtcaga | agatgggaca | tttctgggtc | 600 |

| aactgctacc | tatgctgcaa | agtctcgaag | tacttgatct | ttccattaac | agagacattg | 660 |
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| agttacattc | atgtggatta | tcacaaaaga | gtgtcaaaat | attggatgct | gcttttaggt | 780 |
| atttgggtga | gctgaggaaa | ttagatcttt | cctgcaataa | ggatctaggt | ggaggttttg | 840 |
| aagactcgcc | ggctcagttg | gtcatgctaa | agcatctaca | agtcctagat | cttcaccagt | 900 |
| gctcactaac | agcagatgac | gtgatgtcac | tgacccaggt | cattccttta | ctttcaaatc | 960 |
| ttcaagaatt | ggatttatca | gccaacaaaa | agatgggcag | ttcttctgaa | aacttactca | 1020 |
| gcaggctccg | atttttacca | gcattgaagt | cattagttat | caacaactgt | gctttggaga | 1080 |
| gtgagacttt | tacagctctt | gctgaagcct | ctgttcacct | ctctgctctg | gaagtattca | 1140 |
| acctttcttg | gaacaagtgt | gttggtggca | acttgaagct | gcttctggaa | acactaaagc | 1200 |
| tttccatgtc | tcttcaagtg | ctgaggctga | gcagctgttc | cctggtgaca | gaggatgtgg | 1260 |
| ctctcctggc | atcggtcata | cagacgggtc | atctggccaa | actgcaaaag | ctggacctga | 1320 |
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| aatggtttag | acacttgtta | tatgctgtga | ccaagcttcc | tcagatcact | gagataggaa | 1500 |
| tgaaaagatg | gattctccca | gcttcacagg | aggaagaact | agaatgcttt | gaccaagata | 1560 |
| aaaaaagaag | cattcacttt | gaccatggtg | ggtttcagta | aactgatttc | ccatgtccta | 1620 |
| ctaagctaca | aaccattctc | caaaggaaaa | gaacatgaac | gaattccaga | gtcatgaact | 1680 |
| gaatttcaac | ttctgggcca | tttaatggga | cttatattac | aagagctttg | taaatatata | 1740 |
| tatatattac | atatatatat | gtaatataca | tatatacaca | tatatataat | atacatatat | 1800 |
| aatacacata | tatatgtaaa | tatatata | atatctaata | tgagcatgcc | attattctct | 1860 |
| gtctatgaaa | caaaaatggc | atttttcaat | ggatttgttt | tggatatata | attagttcat | 1920 |
| ttgctgttta | gaageettge | caaaagtgtt | tagattttgg | tactgcaact | gctttcctct | 1980 |
| tgcccagaaa | tgttttgcct | cttcttttcc | tacaagttaa | atgttctaaa | tataaagggg | 2040 |
| tatgtgtgtg | tgtgtgtaat | tctaatgtga | aaggcactag | ctgtctaata | gtttcatgta | 2100 |
| tcattactat | tactatatgt | atcttaatgt | agtctatgta | ggtttttatc | agaaagtgta | 2160 |
| cctttctatg | gtttattatt | ttatattctg | gtgcctttta | tctcagatat | aaaccatgaa | 2220 |
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caaggaagaa aactteetea gaaggagaaa etaageeeca gaetteaaet gteaacaaat
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                                                                     240
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cccaacccaq cqactqqata caqaaqacaq ccacctcaqa gactgctaag cctctcagtt
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Phe Thr Cys Leu Phe Ile Phe 2.0

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Lys Phe Ser Thr Leu Ala Thr Asp Tyr Lys Ser Val Pro Ser Gly Cys

Cys

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Trp Val His Ile Cys Gln Ala Asp Val Tyr His Thr Leu Leu Lys Gly

Phe Lys Ser Val Phe Glu Thr Glu Ser His Val Val Ser Pro Arg Leu 35 40 45

Glu Cys Asn Gln Ser Lys Thr Pro Leu Lys Lys Asn Lys 50 55

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Glu Ser

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<400> 108

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Tyr Lys Lys Gln Asp Ile Leu Pro Gln Leu Arg Ser Asp Lys Ile Thr

Leu Gly Lys Leu Gln Gly Gln Cys Ala Ser Lys Thr Lys Ser Leu Val 40

Ser Ser Leu Thr Ser Tyr Leu Pro Ala Phe Ile Ile Ser Leu Ser

Val Thr Gln Tyr Leu Val Asn Phe Leu Phe Trp His Thr 70

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<212> PRT

<213> Homo sapiens

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Gln Ala His Ser Ile Thr Arg Leu Ser His Ile Gly Val Phe His Phe

Gly Asp Glu Asp Glu Gly Glu Ser Gly Arg Glu

<210> 110

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Glu Asn Ser Asp Gly Asp Leu Ser Lys Val Ile Leu Pro Lys His His 2.0 25

Leu Ala Ile Met Ile Pro Pro Asn Leu Ser Gln Phe Val Tyr Phe Ile 40

Ser Arg Gly Ser Phe Ser Val Leu Ala Ser Cys Val Phe Val Phe Phe

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Arg Ile Ser Leu Leu Lys Glu Ala Gly Gly Thr 85 90

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Ser Leu Trp Arg His Asn Pro Asn Cys Glu Leu Leu Asn 35 40

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Phe Ser Gln Ser Phe Leu Leu Ala Phe Leu Ser Asn Arg Val Leu Leu
    35
               40
Thr Pro Tyr Ile Pro Phe Trp Leu Val Arg Val Ser Phe Ser Ser Ser
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Xaa Asn Asn Lys Leu Leu Gln Leu Phe
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<211> 57
<212> PRT
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10

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Pro Ser Val Lys Trp Lys Arg Glu Glu 50 55

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Gly Met Ala Leu Gly Leu Gly Leu Val Gly Thr Ala Ala Thr Arg Gly 50 55 60

Gly Ser Ser Ala Trp Pro Asp Ser Thr Cys Asn Val Gly Arg Gln Trp 65 70 75 80

Ala Pro Pro Gly Gly Arg Asn Thr Val Arg Ser Met Gln Arg Ala Gly 85 90 95

Asp His Gly Ala Cys Asp Leu Arg Ala His Pro Gly Gln Thr Trp Val

Arg Gly Gly Leu Gly Arg Gln Asp Ser Glu Gly Leu Gln Gly Val Phe
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Val Leu Cys Pro Tyr Thr Gly Asp Leu His Gly Arg Val Arg Ser Ile 130 135 140

Arg Met Leu 145

<210> 116

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Gly Gly Lys Leu Glu Ala His Pro His Phe Thr Gln Thr Leu Gly Ile 50 55

Pro Ile Ser Gly Asn Arg Gly Val Phe

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Asn Ser Arg Leu Arg Ile Ser Lys Asn Ser Pro Arg Ala Xaa Phe Arg 4.0

<210> 118

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Gln Thr Lys His Thr Gly 35

<210> 119

<211> 55

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Asn Val Leu Glu Ser Leu Lys Phe Ile Phe Lys Cys Glu Ser Leu Leu 20 25 30

Phe Val Trp Gly Glu Glu Cys Gln Val Gly Ile Met Asn Gln Ala Leu 35 40 45

Pro Tyr Gln Val Leu Leu Tyr 50 55

<210> 120

<211> 92

<212> PRT

<213> Homo sapiens

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Glu Ser His Thr Leu Gln Val Ile Leu Gly Cys Glu Met Gln Glu Asp 1 5 10 15

Asn Ser Thr Glu Gly Tyr Trp Lys Tyr Gly Tyr Asp Gly Gln Asp His 20 25 30

Leu Glu Phe Cys Pro Asp Thr Leu Asp Trp Arg Ala Ala Glu Pro Arg 35 40 45

Ala Trp Pro Thr Lys Leu Glu Trp Glu Arg His Lys Ile Arg Ala Arg 50 55 60

Gln Asn Arg Ala Tyr Leu Glu Arg Asp Cys Pro Ala Gln Leu Gln Gln 65 70 75 80

Leu Leu Glu Leu Gly Arg Gly Val Leu Asp Gln Gln 85 90

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Met Ile Lys Val Ser Leu Thr Ser Ala Pro Lys Val Ser Ser Leu Glu
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Gly Thr Asn Arg Arg Glu His Ser Asp Thr Gln Gly Pro Leu Ser Val
Pro Trp Lys Pro Ser Asp Leu Cys Arg Pro Ile Ser Val Arg Lys Trp
55
Xaa Xaa Xaa Xaa Xaa Xaa Xaa Arg Thr Thr Gln Ser Ser Trp Gln
Ile Leu Asn Lys Gly
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<223> X=any amino acid
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Met Gly Gly Ala Trp Ser Ile Ala Gly Pro Leu Thr Gly Phe Xaa Phe
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Arg Leu Thr Phe 20

<210> 123

<211> 103

<212> PRT

<213> Homo sapiens

<400> 123

Phe Tyr Phe Leu Phe Ser Phe Val Leu Arg Trp Ser Phe Thr Leu Val 10

69

Thr Gln Ala Gly Val Gln Trp Cys Asp Leu Gly Ser Leu Gln Pro Pro 25

Pro Pro Arg Leu Lys Ala Phe Ser Cys Leu Gly Leu Pro Ser Ser Trp 40

Asp Tyr Arg His Ala Leu Gln Arg Pro Ala Asn Phe Ala Phe Leu Val 50 55

Glu Ile Gly Phe His His Val Gly Gln Ala Gly Pro Gln Leu Leu Thr 70 75

Ser Gly Asp Pro Ser Ile Leu Ala Ser Gln Ser Ala Gly Ile Thr Gly 85 90

Val Thr Ala Val Pro Gly Pro 100

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<223> X=any amino acid

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Met Val Val Ile Gln Ala Xaa Glu Glu Glu Lys Thr Xaa Xaa Xaa Xaa 5 10

2.0 25 30

<210> 125

<211> 95 <212> PRT <213> Homo sapiens

<400> 125

Met Ser Ser Tyr Met Ile Asn Lys Phe Leu Pro Ile Lys Lys Val Lys 10

Ile Pro Gly His Lys Val Phe Ser Thr Asp Ile Met Phe Leu Lys Phe

Val Ser Ile Ala Thr Leu Leu Arg Arg His Thr Asp Ile Ser Glu Asp 40

Leu Arg Val Leu Gln Asn Thr Glu Lys Ile Ser Arg Arg Lys Gly Lys 55

Gly Glu Thr Lys Lys Leu Lys Glu Gly Leu Thr Tyr Lys Trp Asn Asp 7.0

Leu Lys Arg Asn Gly Glu Pro Gly Glu Thr Gly Val Ser Gln Ser

<210> 126

<211> 48 <212> PRT <213> Homo sapiens

<400> 126

Met Ile Lys Tyr Phe Lys Ser Asn Asn Tyr Lys Phe Asn Tyr Tyr Lys

Thr Ser Ser Leu Thr Ser Asp Cys Phe Val Leu Ser Phe Lys Ile Ile

Met Val Cys Leu Arg Val Cys Leu Leu Asn Thr Phe Ala Tyr Leu Pro 40

<210> 127

<211> 98

<212> PRT

<213> Homo sapiens

<400> 127

Met Glu Phe Arg Ser Val Ala Gl
n Val Gly Val Gl
n Trp Arg Asp Leu 1 5 10 15

Gly Leu Gln Pro Leu Pro Leu Gln Phe Lys Gln Phe Tyr Cys Leu $20 \ 25 \ 30$

Ser Leu Ser Ser Ser Trp Asp Tyr Arg His Ser Pro Pro His Pro Ala 35 40 45

Asn Phe Leu Tyr Phe Ala Lys Ile Leu Tyr Ile Ala Lys Arg Phe His 50 55 60

His Val Gly Gln Ala Gly Leu Ala Leu Leu Thr Ser Gly Asp Pro Pro 65 70 75 80

Thr Ser Ala Ser Gln Ser Ala Gly Ile Thr Gly Leu Ser His Cys Ala 85 90 95

Gln Pro

<210> 128

<211> 50

<212> PRT

<213> Homo sapiens

<400> 128

Met Gly Lys Arg Arg Asp Ser Trp Thr Asn Arg Glu Arg Gln Leu Glu 1 5 10 15

Asn Lys Ser Met Gln Lys Ile Ile Tyr Asn Lys Ile Met His Leu Thr 20 25 30

Leu Val Thr Lys Gln Ile Ser Tyr Pro His Phe Ser Leu Ser Val Phe 35 40 45

Val Ser

<210> 129

<211> 16

<212> PRT

<213> Homo sapiens

<400> 129

Met Leu Leu Phe Val Leu Ser Leu Val Phe Gln Tyr Gln Phe Asn Thr 1 5 10 15

<210> 130

<211> 54

<212> PRT

<213> Homo sapiens

<400> 130

Met Ala Leu His Cys Phe Thr Ser Gly Leu Trp Ile Ala Ser Val Arg

Lys Lys Val Lys Met Lys Glu Lys Val Glu Gln Ile Leu Ala Thr Glu 20 25 30

Pro Pro Glu Asp Ser Cys Pro Phe Ser Asn Lys Leu Ser Gly Lys Cys 35 40

Cys Cys His Gly Ser Thr 50

<210> 131 <211> 41 <212> PRT <213> Homo sapiens

<400> 131

Met Cys Ala His Lys Gly Lys Ala Met Arg Glu Arg Thr Gln Pro Glu 10

Gly Gly His Leu Ala Ser Gln Gly Glu Ala Leu Arg Glu Thr Lys Pro 20 25

Ala Arg Leu Gly Thr Val Ala His Gly 35

<210> 132

<211> 35 <212> PRT <213> Homo sapiens

<400> 132

Met Ala Leu Ile Leu Glu Ala Leu Cys Phe Gly Leu Ile Ile Cys 10

Met Asn Arg Glu Ser Ile Ser Thr Leu Ile Phe Tyr Lys His Trp Met

Ser Ile Leu

<210> 133

<211> 58 <212> PRT <213> Homo sapiens

<400> 133

Met Phe Asn Ala Tyr Leu Leu Tyr Asn Asn Gln Val Ile Thr Val Gln 1 5 10 15

73

Ile Lys Gly Pro Lys Cys Phe Arg Tyr Asp Ile Ile Leu Ser Ile Val 20 25 30

Asn Trp Thr Lys Glu Thr Leu Tyr Val Gln Gly Ser Val Glu Gln Pro 40 45 35

Trp Cys Ser Trp Asp Met Leu Pro Arg Cys 50 55

<210> 134 <211> 27 <212> PRT <213> Homo sapiens

<400> 134

Met Met Lys Leu Cys Phe Thr Ala Ser Leu Leu His Gly Ala Leu Leu 1 5 10 15

Trp His Leu Ala Thr Thr Asn Ser Leu Ile Pro 20 25

<210> 135

<211> 46

<212> PRT

<213> Homo sapiens

<400> 135

Met Glu Leu Pro Ser Met Cys Pro Ile Leu Phe Phe Val Thr Val Phe 10

Phe Met Tyr His Thr Pro Ser Cys Pro Ser Ser Val Pro Gln Thr His 20 25 30

Gln Ser His Phe Leu Leu Thr Ala Leu Gly Leu Ala Leu Thr 35 40

<210> 136

<211> 77

<212> PRT

<213> Homo sapiens

<400> 136

Met Thr Cys Pro Gly Gly Glu Thr Gly Trp Gly Cys Leu Arg Met Asp 10

Pro Arg Glu Trp Val Ser Ser Pro Asp Gln Gln Asn Leu Arg Met Cys 25

Ala Trp Ile Gln Pro His Leu Lys Leu Gly Leu His Phe Val Ser Gly 35 40 45

Ala Pro Asn Ala Leu Cys Leu Gly Cys Leu Tyr Ser Trp His Thr Gly 50 55 60

Glu Ala Leu Ser Pro Ala Gly Pro Gly Cys Cys Ser 70

<210> 137 <211> 37 <212> PRT <213> Homo sapiens

<400> 137

Met Glu Gln Glu Ser Val Pro Ser Met Ser Leu Phe Thr Arg Ile Leu 10 15

Ser Gln Pro Ser Leu Phe Pro Trp Gln Ala Leu His Arg Glu Thr Gly 20 25

Lys Arg Ser Thr Val 35

<210> 138

<211> 59

<212> PRT

<213> Homo sapiens

<400> 138

Met Leu Leu Pro Leu Pro Ala Ile Ser Phe Pro Cys Asn Ser Leu Phe 10

His Pro Ala Asp Ala Ser Ser Leu Ser Trp Leu Ser Ser Lys Ser Tyr 20 25

Pro Leu Gly Lys Leu Thr Arg Met Leu Gln Ser Asp Gly Val Ser Pro 45 35 40

Pro Gly Pro Pro Gln Thr Leu Tyr Phe Leu Leu 55

<210> 139

<211> 50

<212> PRT

<213> Homo sapiens

<400> 139

Met Asp Asn Lys Cys Leu Thr Leu Thr Asn Tyr Leu Ala Ile Met Gly

Phe Phe Asp Gln Lys Ser Ser Lys Arg Val Trp Trp Gly Leu Arg Asp 25 30 20

Pro Ser Ser Leu Pro Lys Asn Met Lys Ser Phe His Phe Gln Tyr Val 40

Lys Thr 50

<210> 140

<211> 72

<212> PRT

<213> Homo sapiens

<400> 140

Met Arg Val Val Phe Lys Ile Thr Phe Cys Arg Val Val Cys Ser Thr 5 10 15

Leu Met Leu Lys Gly Ser His Leu Pro Gln Pro Ile Lys Leu Cys Cys

Leu Cys Ser Ala Phe Tyr His Lys Asn Met Thr Phe Lys His Lys Asn 35 40

Thr Leu Tyr Ser Thr Thr Lys Asn Arg Asn Asp Ile Tyr Leu His Cys 55 5.0

Phe Pro Ile Ser Leu His Leu Tyr

<210> 141 <211> 863 <212> PRT <213> Homo sapiens

<400> 141

Met Pro Glu Gln His Lys Asp Pro Arg Val Gln Glu Asn Pro Asp Asp 1 5 10 15 15 Gln Arg Thr Val Pro Glu Val Thr Gly Asp Ala Arg Ser Ala Phe Trp 25 30

Pro Leu Arg Asp Asn Gly Gly Pro Ser Pro Phe Val Pro Arg Pro Gly 35 40 45

Pro Leu Gln Thr Asp Leu His Ala Gln Ser Ser Glu Ile Arg Tyr Asn 50 55 60

His Thr Ser Gln Thr Ser Trp Thr Ser Ser Ser Thr Lys Arg Asn Ala 65 70 75 80

Arg Arg Gly Pro Ala Ser Ser Arg Cys Gln Leu Thr Leu Ser Tyr Ser 100 105 110

Lys Thr Val Ser Glu Asp Arg Pro Gln Ala Val Ser Ser Gly His Thr 115 120 125

Arg Cys Glu Lys Gly Ala Asp Thr Ser Pro Gly Gln Thr Ile Ala Pro 130 \$135\$

Thr Gly Gly Ser Pro Arg Ser His Asp Ser Arg Pro Arg Arg Lys 145 150 150 160

Ile Pro Leu Leu Pro Arg Arg Gly Glu Pro Leu Met Leu Pro Pro 165 170 175

Pro Leu Glu Leu Gly Tyr Arg Val Thr Ala Glu Asp Leu His Leu Glu 180 185 190

Lys Glu Thr Ala Phe Gln Arg Ile Asn Ser Ala Leu His Val Glu Asp 195 200 205

Lys Ala Ile Pro Asp Cys Arg Pro Ser Arg Pro Ser His Thr Leu Ser 210 220

Ser Leu Ala Thr Gly Ala Ser Gly Gly Pro Pro Val Ser Lys Ala Pro 225 230 235 240

Thr Met Asp Ala Gln Gln Asp Arg Pro Lys Ser Gln Asp Cys Leu Gly 245 250 255

Leu Val Ala Pro Leu Ala Ser Ala Ala Glu Val Pro Ala Thr Ala Pro 260 265 270

Val Ser Gly Lys Lys His Arg Pro Pro Gly Pro Leu Phe Ser Ser Ser 275 280 285

Asp Pro Leu Pro Ala Asn Ser Ser His Ser Arg Asp Ser Ala Gln Val 290 295 300

Thr Ser Met Ile Pro Ala Pro Phe Thr Ala Ala Ser Arg Asp Ala Gly 305 310 315 320

Pro Ser Thr Leu Asn Pro Thr Ser Gly Ser Leu Leu Asn Ala Val Asp 340 345 350

Gly Gly Pro Ser His Phe Leu Ala Ser Ala Thr Ala Ala Ala Arg Ala 355 360 365

Gln Arg Ser Glu Val Arg Tyr Asn Gln Arg Ser Gln Thr Ser Arg Thr 370 375 380

Arg Ser Cys Leu Lys Arg Asn Ala Ser Ser Ser Ser His Ser Ser Thr 385 390 395

Glu Gly Leu Gln Glu Val Lys Arg Arg Gly Pro Ala Ser Ser His
405 410 415

Cys Gln Leu Ala His Ser Ser Ser Asn Thr Val Ser Glu Asp Gly Pro 420 425 430

Gln Ala Val Ser Ser Gly His Arg Cys Glu Asn Lys Ala Gly Thr Ala 435 440 445

Pro Gly Gln Thr Leu Ala Pro Arg Gly Gly Ser Pro Arg Ser Gln Ala 450 455 460

Ser Arg Pro His Ile Asn Thr Ala Leu His Val Glu Asp Lys Ala Ile 465 470 475 480

Ser Asp Cys Arg Pro Ser Arg Pro Ser His Thr Leu Ser Ser Leu Ala

485 490 495

Thr Gly Ala Ser Gly Gly Pro Pro Val Ser Lys Ala Pro Thr Met Asp 505 Ala Gln Gln Asp Arg Pro Lys Ser Gln Asp Ser Leu Gly Leu Leu Ala 515 520 Pro Leu Ala Ser Ala Ala Glu Val Pro Ser Thr Ala Pro Val Ser Gly 535 Lys Lys His Arg Pro Pro Gly Pro Leu Phe Ser Ser Asp Pro Leu 550 555 Pro Ala Thr Ser Tyr His Ser Arg Asp Thr Ala Gln Val Thr Ser Leu 565 570 Ile Pro Ala Thr Phe Thr Ala Ala Ser Arg Asp Ala Gly Met Arg Arg 585 Thr Arg Ser Ala Pro Ala Ala Ala Thr Ala Ala Pro Pro Pro Ser Thr 600 Leu Asn Asn Thr Ser Gly Ser Leu Leu Asn Ala Val Asp Gly Pro 615 Ser His Phe Leu Ala Ser Ala Thr Ala Ala Ala Arg Ala Gln Arg Ser 635 625 630 Glu Val Arg Tyr Asn Gln Arg Ser Gln Thr Ser Arg Thr Arg Ser Cys 645 Leu Lys Arg Asn Ala Ser Ser Ser Ser Ser His Ser Ser Thr Glu 660 665 670 Gly Leu Gln Glu Val Lys Arg Arg Gly Pro Ala Ser Ser His Cys 675 680 Gln Leu Ala His Ser Ser Ser Asn Thr Val Ser Glu Asp Gly Pro Gln 695 Ala Val Ser Ser Gly His Arg Cys Glu Asn Lys Ala Gly Thr Ala Pro 705 710 Gly Gln Thr Leu Ala Pro Arg Gly Gly Ser Pro Arg Ser Gln Ala Ser 725

Arg Pro His Ile Asn Ser Ala Leu His Val Glu Asp Lys Ala Ile Ser

Asp Cys Arg Pro Ser Arg Pro Ser His Thr Leu Ser Ser Leu Ala Thr 760

Gly Ala Ser Gly Gly Pro Pro Val Ser Lys Ala Pro Thr Met Asp Ala 775

Gln Gln Asp Arg Pro Lys Ser Gln Asp Cys Leu Gly Leu Leu Ala Pro 785 790 795

Leu Ala Ser Ala Ala Glu Val Phe Ser Thr Ala Pro Val Ser Gly Lys 805 815 810

Lys His Arg Pro Pro Gly Pro Leu Phe Ser Ser Asp Pro Leu Pro 820 825 830

Ala Thr Ser Ser His Ser Gly Asp Ser Ala Gln Asp Thr Ser Leu Ile 835 840 845

Pro Ala Pro Phe Thr Pro Ala Ser Arg Asp Ala Gly Ile Arg Arg 855

<210> 142 <211> 29 <212> PRT <213> Homo sapiens

<400> 142

Met Ser Tyr Leu Ser Leu Leu Leu Ile Ser Ile Phe Met Val Cys Tyr 1 5 10 15

Phe Lys Arg Asn Ser Phe Pro Ile Thr Ile Leu Phe Ser 20 25

<210> 143

<211> 32

<212> PRT

<213> Homo sapiens

<400> 143

Met Pro Trp Pro Met Pro Ile Cys Thr Gly Thr Gln Gly Val Leu Thr 1 5 10

His Arg Gln Gly Pro Pro Pro Ala Ala Val Gly Val Ser Pro His Thr

<210> 144 <211> 29 <212> PRT <213> Homo sapiens

<400> 144

Met Asn Ala Phe Leu Leu Glu Arg Met Thr Glu Ser Gln Ala Met Asp 1 5

Ile Gln Thr Cys Ile Phe Gln Thr Leu Leu Glu Asn Lys 20 25

<210> 145

<211> 48

<212> PRT

<213> Homo sapiens

<400> 145

Met Ile Val Thr Asn Thr Ile Leu Lys Phe Ile His Lys Lys Pro Thr 1.0

Thr Ile Thr Pro Thr Lys Gln His Gly Ile Ile Phe Ser Val Pro Glu 25

Ala Lys Val Arg Ala Leu Leu Cys Phe Leu Leu Arg Met Pro Ser Pro 35 40

<210> 146

<211> 55

<212> PRT

<213> Homo sapiens

<400> 146

Gly Gln Ala Leu Trp Leu Met Pro Val Ile Pro Val Val Ala Lys Ala

Glu Gly Lys Asp His Leu Arg Pro Gly Val Ala Asn Gln Pro Gly Gln

His Ser Lys Thr Leu Phe Leu Gln Lys Lys Asn Phe Ala Lys Leu Ala 40

Glu His Gly Gly Ala Cys Leu 55 5.0

<210> 147

<211> 55

<212> PRT

<213> Homo sapiens

<400> 147

Met Ser Arg Phe Arg Ile Gln Thr Ser Glu Thr Ala Pro Ile Pro Leu 10

81

Val Ser His Pro His Thr Pro Leu Ser Asn Asn Asn Leu His Leu 20 25

Gly Asn Val Cys Tyr Val Pro Gly His Thr Gly Ile Ile Ser Cys Thr 35 40

Pro His Arg His Leu Ile Lys 50 55

<210> 148

<211> 50 <212> PRT <213> Homo sapiens

<400> 148

Met Gln Gly Leu His Leu Pro Gln Gly Leu Gly Thr Cys Tyr Ser Ile

Cys Leu Gln Cys Leu Ser Pro His Gly Tyr Phe Phe Val Ala Val Gly 20 25 30

Leu Ser Ser Asn Val Met Ser Pro Thr Ser Leu Pro Lys Ala Val Leu 35 40 45

Pro Thr 50

<210> 149 <211> 31 <212> PRT <213> Homo sapiens

<400> 149

Met Leu Pro Val Asn Ile Ser His Pro Leu Ser Arg Gly Asn Pro Leu 1 5 10 15

Leu Ser Ser Lys Phe Ser Lys Phe Phe Leu Ile Glu Phe Ser Gln 20 25

<210> 150

<211> 36

<212> PRT

<213> Homo sapiens

<400> 150

Met Asp Tyr Ser Leu Ser Phe Asp Asn Tyr Thr Trp Gly Phe Gly Glu 1 5 10 15

82

Pro Arg Ile Lys Val Gln Ser Phe Asn Asp Leu Leu Ala Pro Gly Leu 20 25 30

Thr Gln Glu His 35

<210> 151

<211> 85

<212> PRT

<213> Homo sapiens

<400> 151

Met Ile Arg Ser Lys Gly Thr Asn Phe Gln Ile Leu Ala Glu Leu Phe 1 5 10 15

Lys Gly Met Asp Phe Leu Trp Leu Gln Leu Ala Arg Leu Phe Gln Lys 20 25 30

Ala Cys Pro Trp Leu Thr Ala Cys Leu Ala Gln Phe Leu Arg Ser Pro $35 \hspace{1cm} 40 \hspace{1cm} 45$

Leu Val Met Glu Asn Arg Ala Asp Arg Ile Gln Met Ala Arg Phe His 50 55 60

Arg Gly Gln Gly Pro Gln Ser Ala Asn Gln Gly Arg Leu Arg Pro 65 70 75 80

Glu Lys Gly Ile Ser

<210> 152

<211> 73

<212> PRT

<213> Homo sapiens

<400> 152

Met Asp Arg Phe Leu Asn Ser Lys Ala Arg Arg Leu Gly Ser Cys Ser 1 5 10 15

His Pro Ala Phe Tyr Leu Leu Cys Val Pro Asp Glu Asp Thr Ser Cys

Ser Thr Met Tyr Leu Pro Leu Lys Arg Ala Asp Pro Asp Gln Leu

Phe Ser Asp Leu Leu Gly Gly Thr Gln Arg Leu Trp Arg Leu Trp Pro 55

Ser Leu Ala Ser Val Glu Ser Gly Leu

<210> 153 <211> 63

<212> PRT

<213> Homo sapiens

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<223> X=any amino acid

<400> 153

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35 40

Met Lys Gln Glu Leu Ser Trp Thr Ile Tyr Asn Leu Leu Arg Tyr 55

<210> 154

<211> 46 <212> PRT

<213> Homo sapiens

<400> 154

Met Arg Cys Leu Leu Ala Asp Ser Ser Leu Gln Met Gln Pro Gly Asp

Val Thr Leu Arg Leu Glu Ser Cys Gly Ser Asn Pro Arg Gln Arg Gln 20 25

Leu His Gln Val Leu Val Trp Val Arg Asn Arg Gly Lys Gly 40

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<210> 155
<211> 72
<212> PRT
<213> Homo sapiens
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<222> (22)..(22)
<223> X=any amino acid
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Met Pro Pro Arg Gly Trp Ala Cys Pro Ser Ser Gly Pro Pro Ala Pro
Gly Pro Gly Arg Trp Xaa Arg Ala Ala Ala Gly Gly Leu Arg Arg Thr
Arg Cys Asp Trp Leu Pro Leu Arg Arg Thr Gln Met Ser Leu Arg Arg
                                40
Ile Asp Leu Leu Pro Ser Pro Ala Gly Gln Ala Gln Ala Gly Ser Glu
Asn Tyr Leu Pro Leu Phe Ile Ser
<210> 156
<211> 20
<212> PRT
<213> Homo sapiens
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<223> X=any amino acid
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<222> (16)..(16)
<223> X=any amino acid
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<223> X=any amino acid

<400> 156

Met Val Phe Ile Phe Ser Thr Thr Ile Xaa Phe Phe Xaa Xaa Glu Xaa 10

Glu Ser Cys Xaa

<210> 157

<211> 66 <212> PRT <213> Homo sapiens

<400> 157

Met Ser Leu Thr Tyr Ser Trp Lys Lys Ser Lys Val Thr Lys Phe Asn

Leu Ser Thr Leu Arg Met Thr Val Thr Asn Lys Asn Arg Thr Val Gln

Lys Cys Ala Lys Asp Thr Arg Lys Leu Asn Asn Ile Asn Ser Met Ile 35 40

Ile Val Ile Leu Tyr Thr Met Glu Ser Lys Gln Ile Phe Phe His Gly

Asn Ser 65

<210> 158

<211> 41

<212> PRT

<213> Homo sapiens

<400> 158

Met Met Thr Gly Glu Ala Arg Glu Ser Gln Ile Ala Leu Tyr Lys Gln 5 10

Arg Phe Arg Glu Phe Arg Glu Glu Gly Arg Thr Ile Tyr Lys Gly Arg

Trp Lys Arg Ser His Leu Ala Glu Gly

<210> 159

<211> 31

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86
<212> PRT
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<223> X=any amino acid
<400> 159
Met Leu Glu Leu Gly Leu Xaa Pro Lys Leu Thr Ser Glu Tyr Arg Phe
                                     10
Pro Pro Asn Cys Met Ile Leu His Ile Trp Ser Gln Leu Glu Val
                                  25
<210> 160
<211> 75
<212> PRT
<213> Homo sapiens
<400> 160
Met Tyr Ile Tyr Ile Cys His His Phe Lys Asn Gln Ala Phe Lys Val
                                      10
Lys Leu Ser Phe Pro His Ile Phe Phe His Ser Ile Phe Tyr Gln Tyr
Arg His Ser Leu Leu Leu Ser Phe Gln Phe Pro Ile Ile Ser Ser
His Pro Ile Phe Cys Ala Ser Ser Val Phe Lys Thr His Ser Pro Ser
    50
                         55
Ala Ala Met Ala Pro Thr Ile Ile Phe Ile Thr
                    70
<210> 161
<211> 36
<212> PRT
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<222> (7)..(13) <223> X=any amino acid

<400> 161

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Met Lys Arg Gly Asn Leu Xaa Xaa Xaa Xaa Xaa Xaa Xaa Gly Thr Pro 1 $$ 5 $$ 10 $$ 15

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Cys Lys Asp Trp Ser His Thr Ala Met Ser Gln Glu Pro Pro Val Leu
            20
                                    25
Val Arg Val Leu
        35
<210> 162
<211> 24
<212> PRT
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<223> X=any amino acid
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<222> (20)..(20)
<223> X=any amino acid
<400> 162
Met Trp Ala Ala Trp Arg Arg Xaa Asn Gly Phe Phe Pro Arg Ile
Pro Gly Lys Xaa Arg Gly Pro Asn
          20
<210> 163
<211> 31
<212> PRT
<213> Homo sapiens
<400> 163
Met Cys His Ser Leu Tyr Arg Phe Leu Asn Cys His Ser Arg Tyr Tyr
Ile Val Tyr Thr Tyr Leu Thr Ile Phe Tyr Trp Cys His His Phe
            20
                                    25
<210> 164
<211> 134
<212> PRT
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Met Ser Lys Val Asp Leu Phe Ile Thr Asp Ser Phe Lys Lys Phe Asn

10

88

Gln Tyr Leu Leu Ala Thr Tyr Ser Thr Ser Gly Thr Gln Gly Ile Trp 25

Ser Thr Thr Met Lys Lys Arg Asp Trp Thr Leu Lys Glu His Arg Ser 40

Cys His Phe 50

<210> 166

<211> 60 <211> 60 <212> PRT <213> Homo sapiens

<400> 166

Met Ser Asp Ser Arg Leu Cys Ser Cys Phe Leu His Thr Leu Ile Phe

Leu Asn Ile Ser Lys Ile Gln Ser Gly Ser Lys Ile Thr Cys Lys Asn

Ile Leu Ala Gln Glu Phe Asp Arg Leu Lys Ile Asn Tyr Leu Lys Tyr 40

Ile Lys Gln Glu Val Tyr Leu Leu Tyr Ser Met Tyr 50 55

<210> 167 <211> 15 <212> PRT

<213> Homo sapiens

<400> 167

Met Val Phe Gln Lys Thr Ser Leu Tyr Ser Asn Asn Ile Leu Leu 10

<210> 168

<211> 106

<212> PRT

<213> Homo sapiens

<400> 168

Cys Pro Ala Ala Tyr Thr Val Phe Leu Thr Arg Ile Ile Val Lys Tyr

Tyr Leu Asn Arg Gly Leu Phe Ser Glu Thr Pro Ser Asn Leu Lys Val 20 25

Glu Glu Lys Gly Trp Val Trp Trp Leu Met Pro Val Thr Pro Ala Leu 40

Trp Glu Ala Glu Ala Gly Gly Ser Leu Glu Leu Ser Leu Arg Pro Gly 55

Trp Ala Thr Pro Ser Leu Pro Lys Asn Thr Lys Met Ser Gln Ala Trp 70 75

Trp Cys Thr Pro Val Val Pro Ala Ala Leu Gly Ala Glu Val Gly Gly 90

Arg Leu Gly Pro Arg Arg Trp Arg Leu Gln 100

<210> 169

<211> 19

<212> PRT <213> Homo sapiens

<400> 169

Met Gly Pro Asp Arg Leu Lys Gln Lys Ser Asn Thr Ala Val Val Ser 10

Arg Trp Ile

<210> 170 <211> 47 <212> PRT

<213> Homo sapiens

<220>

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<223> X=any amino acid

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<222> (16)..(16) <223> X=any amino acid

<400> 170

Met Asp Xaa Xaa Lys Trp Arg Met Arg Arg Gln Pro Xaa Ile Asn Xaa

Met Tyr Gln Thr Val Thr Ile Cys Glu Glu Tyr Cys Val Tyr Thr Asn 25

Arg Lys Gln Leu Lys Ala Phe Asn Met Cys Gly Trp Gly Glu Arg

<210> 171 <211> 197 <212> PRT <213> Homo sapiens

<400> 171

Gln Glu Ala Gln Ile Met Lys Lys Leu Arg His Asp Lys Leu Val Pro

Leu Tyr Ala Val Val Ser Glu Glu Pro Ile Tyr Ile Val Thr Glu Phe

Met Ser Lys Gly Ala Tyr Ser Leu Ser Ile Arg Asp Trp Asp Glu Ile 40

Arg Gly Asp Asn Val Lys His Tyr Lys Ile Arg Lys Leu Asp Asn Gly

Gly Tyr Tyr Ile Thr Thr Arg Ala Gln Phe Asp Thr Leu Gln Lys Leu 70 75

Val Lys His Tyr Thr Glu His Ala Asp Gly Leu Cys His Lys Leu Thr 85

Thr Val Cys Pro Thr Val Lys Pro Gln Thr Gln Gly Leu Ala Lys Asp

Ala Trp Glu Ile Pro Arg Glu Ser Leu Arg Leu Glu Val Lys Leu Gly 120

Gln Gly Cys Phe Gly Glu Val Trp Met Gly Thr Trp Asn Gly Thr Thr 135

Lys Val Ala Ile Lys Thr Leu Lys Pro Gly Thr Met Met Pro Glu Ala 145 150 155

Phe Leu Gln Glu Ala Gln Ile Met Lys Lys Leu Arg His Asp Lys Leu

175 165 170

Val Pro Leu Tyr Ala Val Val Ser Glu Glu Pro Ile Tyr Ile Val Thr 185 180

Glu Phe Met Ser Lys 195

<210> 172

<211> 59

<212> PRT

<213> Homo sapiens

<220>

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<223> X=any amino acid

<400> 172

Met Cys Ile Met His Ile Asn Thr Phe Asn Leu Cys Asn His Leu Met 5

Arg Trp Leu Leu Lys Ser Pro Leu Cys Thr Xaa Xaa Xaa Xaa Xaa 25 20

35

Xaa Gln Lys Pro Lys Pro Thr Val His Gly Ile

<210> 173

<211> 56

<212> PRT

<213> Homo sapiens

<220>

<221> MISC_FEATURE

<222> (14)..(21) <223> X=any amino acid

<400> 173

Met Lys Pro Ile Arg Gln Leu Val Pro Phe Thr Leu Glu Xaa Xaa Xaa 5

Xaa Xaa Xaa Xaa Leu Tyr Leu Glu His Leu Thr Cys Arg Lys Arg 20 25

Arg Gly Lys Thr Phe Leu Gly Lys Arg Lys Ala Val Ala Val Pro Lys 40

Ser Lys His Phe Trp Gln Gly Phe 50 55

<210> 174

<211> 104

<212> PRT

<213> Homo sapiens

<400> 174

Met Leu Lys His Leu Gln Val Leu Asp Leu His Gln Cys Ser Leu Thr 10

Ala Asp Asp Val Met Ser Leu Thr Gln Val Ile Pro Leu Leu Ser Asn

Leu Gln Glu Leu Asp Leu Ser Ala Asn Lys Lys Met Gly Ser Ser Ser 40

Glu Asn Leu Leu Ser Arg Leu Arg Phe Leu Pro Ala Leu Lys Ser Leu 55

Val Ile Asn Asn Cys Ala Leu Glu Ser Glu Thr Phe Thr Ala Leu Ala 70

Glu Ala Ser Val His Leu Ser Ala Leu Glu Val Phe Asn Leu Ser Trp 90

Glu Gln Val Cys Trp Trp Ala Thr 100

<210> 175

<211> 490

<212> PRT <213> Homo sapiens

<400> 175

Met Ser Gln Thr Arg Lys Lys Thr Ser Ser Glu Gly Glu Thr Lys Pro

Gln Thr Ser Thr Val Asn Lys Phe Leu Arg Gly Ser Asn Ala Glu Ser

Arg Lys Glu Asp Asn Asp Leu Lys Thr Ser Asp Ser Gln Pro Ser Asp 40 35

- Trp Ile Gln Lys Thr Ala Thr Ser Glu Thr Ala Lys Pro Leu Ser Ser 50 55 60
- Glu Met Glu Trp Arg Ser Ser Met Glu Lys Asn Glu His Phe Leu Gln 65 70 75 80
- Lys Leu Gly Lys Lys Ala Val Asn Lys Cys Leu Asp Leu Asn Asn Cys 85 90 95
- Gly Leu Thr Thr Ala Asp Met Lys Glu Met Gly Glu Ala Phe Glu Met 100 \$105
- Ile Pro Glu Leu Glu Glu Leu Asn Leu Ser Trp Asn Ser Lys Val Gly 115 120 125
- Gly Asn Leu Pro Leu Ile Leu Gln Lys Phe Gln Lys Gly Ser Lys Ile 130 135 140
- Phe Leu Gly Gln Leu Leu Pro Met Leu Gln Ser Leu Glu Val Leu Asp 165 170 175
- Leu Ser Ile Asn Arg Asp Ile Val Gly Ser Leu Asn Ser Ile Ala Gln
 180 185 190
- Gly Leu Lys Ser Thr Ser Asn Leu Lys Val Leu Lys Leu His Ser Cys 195 200 205
- Gly Leu Ser Gln Lys Ser Val Lys Ile Leu Asp Ala Ala Phe Arg Tyr 210 215 220
- Leu Gly Glu Leu Arg Lys Leu Asp Leu Ser Cys Asn Lys Asp Leu Gly 225 230 235 240
- Gly Gly Phe Glu Asp Ser Pro Ala Gln Leu Val Met Leu Lys His Leu 245 250250255
- Gln Val Leu Asp Leu His Gln Cys Ser Leu Thr Ala Asp Asp Val Met 260 265 270
- Ser Leu Thr Gln Val Ile Pro Leu Leu Ser Asn Leu Gln Glu Leu Asp 275 280 285

Leu Ser Ala Asn Lys Lys Met Gly Ser Ser Ser Glu Asn Leu Leu Ser 290 295 300

Arg Leu Arg Phe Leu Pro Ala Leu Lys Ser Leu Val Ile Asn Asn Cys 305 310 310 315

Ala Leu Glu Ser Glu Thr Phe Thr Ala Leu Ala Glu Ala Ser Val His 325 330 335

Leu Ser Ala Leu Glu Val Phe Asn Leu Ser Trp Asn Lys Cys Val Gly 340 345 350

Gly Asn Leu Lys Leu Leu Leu Glu Thr Leu Lys Leu Ser Met Ser Leu 355 360 365

Gln Val Leu Arg Leu Ser Ser Cys Ser Leu Val Thr Glu Asp Val Ala 370 375380

Leu Leu Ala Ser Val Ile Gln Thr Gly His Leu Ala Lys Leu Gln Lys 385 390 395 400

Leu Asp Leu Ser Tyr Asn Asp Ser Ile Cys Asp Ala Gly Trp Thr Met 405 410 415

Phe Cys Gln Asn Val Arg Phe Leu Lys Glu Leu Ile Glu Leu Asp Ile 420 425 430

Ser Leu Arg Pro Ser Asn Phe Arg Asp Cys Gly Gln Trp Phe Arg His 435 440445

Leu Leu Tyr Ala Val Thr Lys Leu Pro Gln Ile Thr Glu Ile Gly Met 450 455 460

Lys Arg Trp Ile Leu Pro Ala Ser Gln Glu Glu Glu Leu Glu Cys Phe 465 470 470 475

Asp Gln Asp Lys Lys Lys Lys His Ser Leu 485 490

<210> 176

<211> 136

<212> PRT

<213> Homo sapiens

<400> 176

Gln Glu Ile Ser Ser Ser Ser Lys Ser Gly Arg Lys Gly Ser Lys Ala $20 \ \ 25 \ \ 30$

Thr Ile Ser Phe Met Ser Ala Val Val Asn Pro Gln Leu Phe Lys Ser 35 40 45

Arg His Leu Leu Thr Ala Phe Leu Pro Ser Phe Cys Arg Lys Cys Ser 50 55 60

Phe Phe Ser Ile Leu Asp Leu His Ser Ile Ser Glu Leu Arg Gly Leu 65 70 75 80

Ala Val Ser Glu Val Ala Val Phe Cys Ile Gln Ser Leu Gly Trp Glu 85 90 95

Ser Leu Val Leu Arg Ser Leu Ser Ser Phe Leu Leu Ser Ala Leu Glu 100 105 110

Pro Leu Arg Asn Leu Leu Thr Val Glu Val Trp Gly Leu Val Ser Pro 115 120 125

Ser Glu Glu Val Phe Phe Leu Val